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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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1: gb_ba:*
2: gb_htg:
3: gb_ln:*
4: qb_om:*
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Pred. No. is the number of results predicted by chance to have

score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

	ed. 1 to 20) 3. modulation of stal3 expression 6159694-A 87 12-DEC-2000; .ocation/Qualifiers	classifi (bases rras,J.C tisense tent: US	REFERENCE AUTHORS TITLE JOURNAL FEATURES
		nknown	WORDS RCE
	GI:14104642	AR121066 AR121066.1	ACCESSION VERSION
linear	20 bp DNA from patent US 6159694.	AR121066 Sequence 87	- C N C
	ALIGNMENTS		
AX6	5 6 AX68914	67.	UT A
ARJ	0 6 AK30/96	67.	<b>A</b> (.
AR3	0 6 AR30796	67.	42
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A25	3 6 A25721	. 8 59.	26
AX/	1 11 DOGINS	8 69.	
AX6	7 6 AX68763	. 8 69.	23
AX6	7 6 AX68763	8 69.	22
BD1	6 6 BD10727	.2 71.	20
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LOCUS	AR121066 20 bp DNA linear PAT 16-MAY-2001
DEFINITION	94.
ACCESSION	AR121066
VERSION	AR121066.1 GI:14104642
KEYWORDS	
SOURCE	Unknown.
ORGANISM	Unknown.
	Unclassified.
REFERENCE	1 (bases 1 to 20)
AUTHORS	Karras, J.G.
TITLE	Antisense modulation of stat3 expression
JOURNAL	Patent: US 6159694-A 87 12-DEC-2000;
FEATURES	Location/Qualifiers

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Sequence 1865 from Patent EP1281758
AX689133
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Antisense modulation of stat3 expression Patent: US 6159694-A 23 12-DEC-2000;
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AR121002
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Sequence 1866 from Patent EP1281758
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/mol_type="genomic DNA"
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/mol_type="genomic DNA"
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Pred. No. 1.4e+04;
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Local Similarity 89.5%;
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                                                                                                                                                                                ch 79.0%; Score 15.8; DB 6; Length 25; l Similarity 89.5%; Pred. No. 1.4e+04; 17; Conservative 0; Mismatches 2; Indels
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Sequence 1868 from Patent EP1281758.
AX689136
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Sequence 1870 from Patent EP1281758
AX689138
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Sequence 1869 from Patent EP1281758.
AX689137
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AX689138.1 GI:29411844
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo
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/mol_type="genomic DNA"
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Sequence 1864 from Patent EP1281758
AX689132
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                               Homo sapiens (human)
Homo sapiens
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                        Ling,V. and Dunussi-Joannopoulos,K. GL50 nucleic acids and uses therefor Patent: US 6521749-A 17 18-FEB-2003;
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                                                                                     Unknown
                                                                                                                                  Sequence 17 from patent US 6521749. AR282672
                                                                                                                                                                                                                                               1 GCTCCAGCATCTGCTGCT 18
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                                                                         Unclassified
                                                                                                                          AK282672.1
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Mammalia; Eutheria; Frimates, Catarrhini; Hominidae; Homo.
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/mol_type-"genomic DNA"
/db_xref="taxon:9606"
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                                                                                                                                                                                                                  Four human zinc-finger-containing proteins :
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Mammalia; Eutheria; Primates;
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                                                     /mol_type="genomic DNA"
/db_xref="taxon:9606"
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/mol_type="genomic DNA"
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Pred No 7 9e+04;
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                           1 GCTCCAGCATCTGCTGCTT 19
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